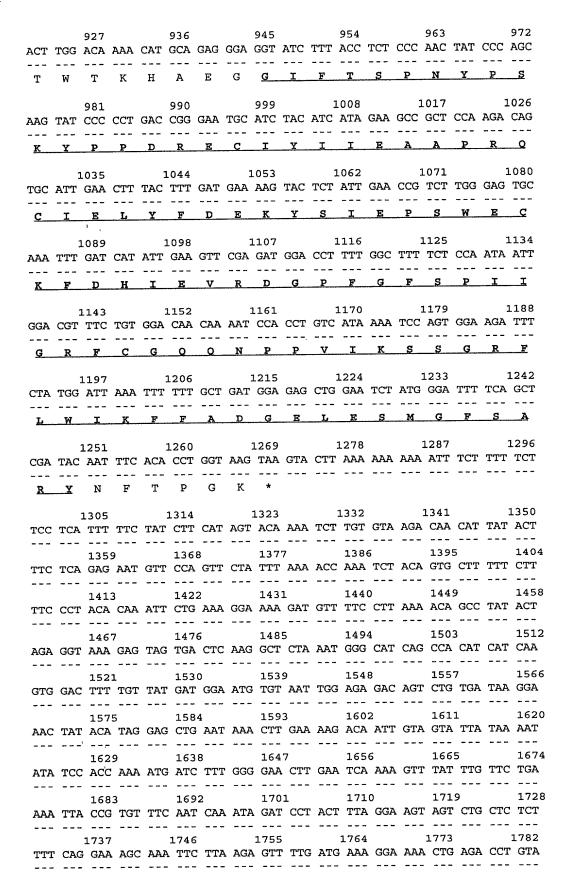
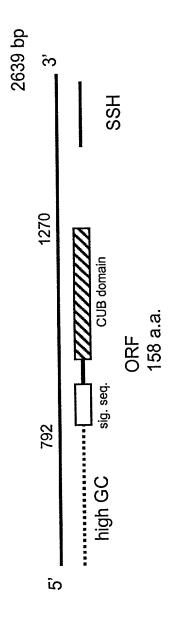
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			225			234			243						261			270
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			333			342									369			378
															CCC			
			387			396			40!			414			423			432
	TCG	GGG	CGT	CCC	GGG										GCG			
															475			
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TTG	AAG	TTT	GTA	TTC	CAT	GTT	AAA	GAA	GTA	ACA	TCT	AAA	IAC	AGC	TTT	GAI	ACI
		216	9		217	8		218	7		2196			220			2214
CAG	TTA	AAA	AAC	TAA	AAT	TTT	AAA	AAT	TAT	TAA	TAT	AAG	TTT	AAT	GAT	GAC	TTT
		2223			2232		:	2241			2250					-	2268
		2223			2232		:	2241			2250					-	
СУД	: ידביד	2223 GAC	ATC	ATG	2232 GGG	TAT	GTT	2241 AAA	TCA	AGT	2250 ATT	TAC	TGT	AGC	ATA	TAT	ATT
СУД	: ידביד	2223 GAC	ATC	ATG	2232 GGG 	TAT	GTT	2241 AAA	TCA	AGT	2250 ATT	TAC	TGT	AGC	ATA	TAT	ATT
CAT	TAT	2223 GAC 	ATC 	ATG	2232 GGG 	TAT 	GTT 	2241 AAA 229	TCA 	AGT	2250 ATT 230	TAC	TGT	AGC 231	ATA 	TAT	ATT 2322
CAT	TAT	2223 GAC 227	ATC 7	ATG	2232 GGG 228 GAA	TAT 6 TGT	GTT 	2241 AAA 229: TAA	TCA 5 TAA	AGT	2250 ATT 230 CAC	TAC	TGT	AGC 231 ATT	ATA 3 GTG	TAT	ATT 2322 TTA
CAT	TAT	2223 GAC 227	ATC 7	ATG	2232 GGG 228 GAA	TAT 6 TGT	GTT 	2241 AAA 229: TAA	TCA 5 TAA	AGT	2250 ATT 230 CAC	TAC 4 TAA	TGT	AGC 231 ATT	ATA 3 GTG	TAT	ATT 2322 TTA
CAT AGC	TAT	2223 GAC 227 AAG 	ATC 7 CAT 	ATG TAG	2232 GGG 228 GAA 234	TAT 6 TGT 	GTT TTT	2241 AAA 229: TAA 234	TCA 5 TAA 	AGT	2250 ATT 230- CAC 235:	TAC 4 TAA 	TGT 	231 ATT 236	ATA 3 GTG 	TAT	ATT 2322 TTA 2376
CAT AGC	TAT	2223 GAC 227 AAG 233	ATC 7 CAT	TAG	2232 GGG 228 GAA 234	TAT 6 TGT 0	GTT	2241 AAA 229: TAA 234 ACT	TCA TAA GAC	AGT TAT	2250 ATT 230- CAC 235- AAT	TAC 1 TAA 8 CAT	TGT AGG 	AGC 231 ATT 236 TAT	ATA 3 GTG 7 TTC	TAT GTT	ATT 2322 TTA 2376 TGC
CAT AGC	TAT	2223 GAC 227 AAG 233	ATC 7 CAT	TAG	2232 GGG 228 GAA 234	TAT 6 TGT 0	GTT	2241 AAA 229: TAA 234 ACT	TCA TAA GAC	AGT TAT	2250 ATT 230- CAC 235- AAT	TAC 1 TAA 8 CAT	TGT AGG 	AGC 231 ATT 236 TAT	ATA 3 GTG 7 TTC	TAT GTT	ATT 2322 TTA 2376 TGC
CAT AGC	TAT	2223 GAC 227 AAG 233 CTT	ATC 7 CAT 1 TGC	TAG	2232 GGG 228 GAA 234 TAA	TAT 6 TGT 0 TGG	GTT TTT	2241 AAA 229! TAA 234 ACT	TCA TAA 9 CAC	AGT TAT	2250 ATT 230 CAC 235 AAT	TAC TAA TAA CAT	TGT AGG 	AGC 231 ATT 236 TAT	ATA GTG 7 TTC	TAT GTT	ATT 2322 TTA 2376 TGC
CAT AGC	TAT TTT ATG	2223 GAC 227 AAG 233 CTT	ATC 7 CAT 1 TGC	TAG	2232 GGG 228 GAA 234 TAA 	TAT 6 TGT 0 TGG	GTT TTT ATT	2241 AAA 2299 TAA 234 ACT 240	TCA TAA 9 CAC	AGT TAT	2250 ATT 230 CAC 235 AAT 241	TAC 4 TAA 8 CAT	TGT AGG 	AGC 231 ATT 236 TAT 	ATA GTG 7 TTC	GTT	ATT 2322 TTA 2376 TGC 2430
CAT AGC ATT	TAT	2223 GAC 227 AAG 233 CTT 238	ATC 7 CAT 1 TGC	TAG	2232 GGG 228 GAA 234 TAA 239	TAT 6 TGT 0 TGG 4	GTT TTT ATT	2241 AAA 229! TAA 234 ACT 240 TCC	TCA TAA TAA CAC	AGT TAT AGA	2250 ATT 230 CAC 235 AAT 241 GTA	TAC TAA TAA CAT CAA	TGT AGG GGG	AGC 231 ATT 236 TAT 242 TTA	ATA 3 GTG 7 TTC 1 GCT	TAT GTT ATG	ATT 2322 TTA 2376 TGC 2430 TTT
CAT AGC ATT	TAT	2223 GAC 227 AAG 233 CTT 238 CGA	ATC 7 CAT 1 TGC	TAG	2232 GGG 228 GAA 234 TAA 239	TAT 6 TGT 0 TGG 4	GTT TTT ATT	2241 AAA 2299 TAA 234 ACT 240 TCC	TCA TAA CAC	AGT TAT AGA	2250 ATT 230 CAC 235 AAT 241 GTA	TAC 1 TAA 8 CAT 2 CAA	TGT AGG GGG	AGC 231 ATT 236 TAT 242 TTA	ATA 3 GTG 7 TTC	TAT GTT ATG	ATT 2322 TTA 2376 TGC 2430 TTT
CAT AGC ATT TAC	TAT TTT ATG	2223 GAC 227 AAG 233 CTT 238 CGA	ATC 7 CAT 1 TGC 5 ACT	TAG	2232 GGG 228 GAA 234 TAA 239	TAT 6 TGT 0 TGG 4 AAG	GTT TTT ATT	2241 AAA 229: TAA 234 ACT 240 TCC 	TCA TAA OCAC CAC CAA	AGT TAT AGA	2250 ATT 230 CAC 235 AAT 241 GTA 246	TAC TAA TAA CAT CAT CAA	TGT AGG GGG	AGC 231 ATT 236 TAT 242 TTA 247	ATA 3 GTG 7 TTC 1 GCT	TAT GTT ATG TAA	ATT 2322 TTA 2376 TGC 2430 TTT 2484
CAT AGC ATT TAC	TAT TTT ATG AGT	2223 GAC 227 AAG 233 CTT 238 CGA	ATC 7 CAT 1 TGC 5 ACT	TAG	2232 GGG 228 GAA 234 TAA 239 TTG 244	TAT 6 TGT 0 TGG 4 AAG	GTT ATT ATT	2241 AAA 2299 TAA 234 ACT 240 TCC 245 AGT	TCA TAA 9 CAC 3 CAA 7 TGT	AGT TAT AGA	2250 ATT 230 CAC 235 AAT 241 GTA 246 TGG	TAC 4 TAA 8 CAT 2 CAA 6 AAA	AGG	231 ATT 236 TAT 242 TTA 247 AGG	ATA 3 GTG 7 TTC 1 GCT	TAT GTT ATG TAA	ATT 2322 TTA 2376 TGC 2430 TTT 2484
CAT AGC ATT TAC	TAT TTT ATG AGT	2223 GAC 227 AAG 233 CTT 238 CGA	ATC 7 CAT 1 TGC 5 ACT	TAG	2232 GGG 228 GAA 234 TAA 239 TTG 244	TAT 6 TGT 0 TGG 4 AAG	GTT ATT ATT	2241 AAA 2299 TAA 234 ACT 240 TCC 245 AGT	TCA TAA CAC TGT	AGA AAA	2250 ATT 230- CAC 235 AAT 241 GTA 246 TGG	TAC TAA TAA CAT CAA CAA AAA	AGG	AGC 231 ATT 236 TAT 242 TTA 247 AGG	ATA GTG TTC 1 GCT	TAT GTT ATG TAA	ATT 2322 TTA 2376 TGC 2430 TTT 2484 TTT
CAT AGC ATT TAC GTT	TAT TTT ATG AGT	2223 GAC 227 AAG 233 CTT 238 CGA TCA	ATC 7 CAT 1 TGC 5 ACT	TAG	2232 GGG 228 GAA 234 TAA 239 TTG 244	TAT G TGT O TGG AAG TGG AAG	GTT TTT ATT TAT	2241 AAA 2299 TAA 234 ACT 240 TCC 245 AGT 	TCA TAA 9 CAC 3 CAA 7 TGT	AGA AAG	2250 ATT 230 CAC 235 AAT 241 GTA 246 TGG 252	TAC TAA TAA CAT CAA AAA	AGG ATG GGT	AGC 231 ATT 236 TAT 242 TTA 247 AGG 252	ATA 3 GTG 7 TTC 1 GCT	TAT GTT ATG TAA TTT	ATT 2322 TTA 2376 TGC 2430 TTT 2484 TTT 2538
CAT AGC ATT TAC GTT	TAT TTT ATG AGT TGT	2223 GAC 227 AAG 233 CTT 238 CGA TCA TCA	ATC 7 CAT 1 TGC 5 ACT	TAGA	2232 GGG 228 GAA 234 TAA 239 TTG	TAT G TGT O TGG AAG TGG AAAG	GTT TTT ATT TAT	2241 AAA 2299 TAA 234 ACT 240 TCC 245 AGT 251 CCT	TCA TAA GAA TGT	AGA AAG AAA	2250 ATT 230 CAC 235 AAT 241 GTA 246 TGG 252 AGA	TAC TAA TAA CAT CAA AAA GAC	TGT AGG GGG ATG	AGC 231 ATT 236 TAT 242 TTA 247 AGG 252 GAC	ATA GTG TTC GCT STAT	TAT GTT ATG TAA TTT CGGG	ATT 2322 TTA 2376 TGC 2430 TTT 2484 TTT 2538 CAT
CAT AGC ATT TAC GTT	TAT TTT ATG AGT TGT	2223 GAC 227 AAG 233 CTT 238 CGA TCA TCA	ATC 7 CAT 1 TGC 5 ACT	TAGA	2232 GGG 228 GAA 234 TAA 239 TTG	TAT G TGT O TGG AAG TGG AAAG	GTT TTT ATT TAT	2241 AAA 2299 TAA 234 ACT 240 TCC 245 AGT 251 CCT	TCA TAA GAA TGT	AGA AAG AAA	2250 ATT 230 CAC 235 AAT 241 GTA 246 TGG 252 AGA	TAC TAA TAA CAT CAA AAA GAC	TGT AGG GGG ATG	AGC 231 ATT 236 TAT 242 TTA 247 AGG 252 GAC	ATA GTG TTC GCT STAT	TAT GTT ATG TAA TTT CGGG	ATT 2322 TTA 2376 TGC 2430 TTT 2484 TTT 2538 CAT
CAT AGC ATT TAC GTT	TAT TTT ATG AGT TGT	22233 GAC 227 AAG 233 CTT 243 TCA 245 TGA	ATC 7 CAT 1 TGC ACT 9 GAT	TAGA	2232 GGG 228 GAA 234 TAA 239 TTG TAG TAG	TAT 6 TGT 0 TGG AAAG 8 TGC	GTT TTT ATT TAT TAG	2241 AAA 229 TAA 234 ACT 240 TCC 245 AGT 	TCA TAA CAC TGT TGT	AGT AGA AAA	2250 ATT 230 CAC 235 AAT 241 GTA 246 TGG 252 AGA	TAC TAA TAA CAT CAA AAA GAC	AGG ATG AGG ATG	231 ATT 236 TAT 242 TTA 247 AGG	ATA GTG TTC GCT TAT	TAT GTT ATG TAA TTT CGGG	ATT 2322 TTA 2376 TGC 2430 TTT 2484 TTT 2538 CAT
CAT AGC ATT TAC GTT CTT	TAT TTTT ATG AGT	22233 GAC 2277 AAG 233 CTT 238 CGA 243 TCA 243 TCA 254	ATC 7 CAT 1 TGC 5 ACT 9 GAT	TAGA TAGA TATT	2232 GGG 228 GAA 234 TAA 239 TTG 250 TGA	TAT 6 TGT 0 TGG 4 AAG 8 TGC 4 AAG 8 ATA	GTT TTT ATT TAT	2241 AAA 2299 TAA 234 ACT 245 AGT 251 CCT 256	TCA TAA O CAC TGT GTA TGTA	AGT TAT AGA AAAG	2250 ATT 230 CAC 235 AAT 241 GTA 252 AGA	TAC 4 TAA 8 CAT 2 CAA 6 AAA 0 GAC	AGG	AGC 231 ATT 236 TAT 242 TTA AGG 252 GAC 2558	ATA 3 GTG 7 TTC 1 GCT 5 TAT	TAT GTT ATG TAA TTT CGGG	ATT 2322 TTA 2376 TGC 2430 TTT 2484 TTT 2538 CAT 2592
CAT AGC ATT TAC GTT CTT	TAT TTTT ATG AGT	22233 GAC 2277 AAG 2333 CTT 238 CGA 243 TCA 245 CGA 254 CGA 255 CGG	ATC 7 CAT 1 TGC 5 ACT 9 GAT	TAGA TAGA TATT	2232 GGG 228 GAA 234 TAA 239 TTG 250 TGA	TAT 6 TGT 0 TGG 4 AAG 8 TGC ATA	GTT TTT ATT TAT TAA	2241 AAA 2299 TAA 234 ACT 245 AGT 256 TTCC	TCA TAA CAA TGT GTA TGT S	AGT TAT AGA AAG CCT CTG	2250 ATT 230 CAC 235 AAT 241 GTA 252 AGA 257 GGT	TAC 4 TAA 8 CAT 2 CAA 6 AAA 0 GAC 4 TTG	AGG	231 ATT 236 TAT 242 TTA 247 AGG 252 GAC 258	ATA GTG TTC TTC TATA STATA ATA ATA ATA	TAT GTT ATG TAA TTT CGGG	2322 TTA 2376 TGC 2430 TTT 2484 TTT 2538 CAT 2592 ATA
CAT AGC ATT TAC GTT CTT	TAT TTTT ATG AGT	22233 GAC 227 AAG 2333 CTT 243 TCA 249 TGA 254	ATC 7 CAT 1 TGC ACT 9 GAT 7 TAA	TAGA TAGA TATT	2232 GGG 228 GAA 234 TAA 239 TTG 250 TGA 255 AGT	TAT6 TGT4 4 AAG8 TGC6 6 TGC6 6 TGC	GTT TTT ATT TAT TAA	2241 AAA 229 TAA 234 ACT 245 AGT 251 CCT 256	TCA TAA OCAC TGT GTA TGT TGT TGTA	AGT TAT AGA AAG CCT CTG	2250 ATT 230 CAC 235 AAT 241 GTA 252 AGA 257 GGT	TAC 4 TAA 8 CAT CAA 6 AAA OGAC 4 TTG	AGG	231 ATT 236 TAT 242 TTA AGG 	ATA GTG TTC GCT TAT SATA ATA ATA ATA ATA ATA	TAT GTT ATG TAA TTT CGGG	2322 TTA 2376 TGC 2430 TTT 2484 TTT 2538 CAT 2592 ATA
CAT AGC ATT TAC GTT CTT GTT	TAT TTTT ATG AGT TGT AACC CTA	22233 GAC 227 AAG 2333 CTT 238 CGA 249 TGA 254 GGT 266	ATC	TAG	2232 GGG 228 GAA 234 TAA 239 TTG TAG TAG AGT AGT 261	TAT 6 TGT 0 TGG 4 AAG 8 TGC 4 AAAG 6 TGC 1 AAAA	TTT ATT TAT TAG TAA	2241 AAA 2299 TAA 234 ACT 245 AGT 251 CCT 256 TTC	TCA TAA TAA CAA TGT GTA TGTA TGTA TGTA TGTA TGTA	AGT TAT AGA AAA CCT CTG	2250 ATT 230 CAC 235 AAT 241 GTA 252 AGA 257 GGT 262	TAC4 CAT6 AAA0 GAC4 TTG8	AGG ATG GGT	231 ATT 236 TAT 242 TTA 247 AGG 252 GAC 258	ATA3 GTG7 TTC1 GCT29 ATAT3 ATAT3 ATAT3 ATAT3 ATAT	TAT GTT ATG TAA TTT CGGG	2322 TTA 2376 TGC 2430 TTT 2484 TTT 2538 CAT 2592 ATA
CAT AGC ATT TAC GTT CTT GTT	TAT TTTT ATG AGT CTA	22233 GAC 2277 AAG 2333 CTT 238 CGA 249 TGA 254 GGI 2564 CGA	ATC	TAGA TATT TATT TATA	2232 GGG 228 GAA 234 TAA 239 TTG TAG TAG AGT AAT	TAT 6 TGT 0 TGG 4 AAG 8 TGC ATA 66 TGC 1 TGC	TTT ATT TAT TAG TAA	2241 AAA 2299 TAA 234 ACT 245 AGT 256 TTC 261	TCA TAA TAA CAA TGT GTA STA ATT ATT ATT ATT A	AGA AAAA CCTG	2250 ATT 230 CAC 235 AAT 241 GTA 252 AGA 257 GGT 262 AAA	TAC 4 TAA 8 CAT 2 CAA 6 AAA 4 TTG 8 AAA	AGG ATG GGT AGT AAAA	231 ATT 236 TAT 242 TTA 247 AGG 252 GAC 258 AAA	ATA 3 GTG 7 TTC 1 GCT 1 S ATA 3 C ATG 3 A AA	TAT GTT ATG TAA TTT CGGG	2322 TTA 2376 TGC 2430 TTT 2484 TTT 2538 CAT 2592 ATA

FIG.



GC content:
5' non-coding 72%
coding 42%
3' non-coding 30%

A
43.5% identity in 115 residues overlap
BPC-1 33 IFTSPNYPSKYPPDRECIYIIEAAPRQCIELYFDEKYSIEPSWECKFDHIEVR CUB domain 75 IFTSPNFPDRYPPNIDCVRVIHSRPQHDVVVKFHHVFHIESTYDKIDAGEECPNDFIEFR ***** * ** * * * * * * * * * * * * * *
BPC-1 86 DGPFGFSPIIGRFCGQQNPP-VIKSSGRFLWIKFFADGELESMGFSARYNFTPGK CUB domain 135 DGRYGFSPLIARFCGDRMPKREIRAVSGFLWIRFRSDSMLEYQGFSAEYAIVPSK ** **** * *** * * **** * * * **** * * *
B
35.3% identity in 102 residues overlap in CUB domain 1 of BMP-1
BPC-1 130 GIFTSPNYPSKYPPDRECIYIIEAAPRQCIELYFDEKYSIEPSWECKFDHIEVRDGPFGF BMP-1 336 GNFSSPEYPNGYSAHMHCVWRISVTPGEKIILNFTSM-DLYRSRLCWYDYVEVRDGFWRK * * * * * * * * * * * * * * * * * * *
BPC-1 190 SPIIGRFCGQQNPPVIKSSGRFLWIKFFADGELESMGFSARY BMP-1 395 VWVRGRFCGGKLPEPIVSTDSRLWVEFRSSSNWVGKGFFAVY ***** * * * * * * * * * * * * * * * *
39.4% identity in 104 residues overlap in CUB domain 2 of BMP-1
BPC-1 128 EGGIFTSPNYPSKYPPDRECIYIIEAAPRQCIELYFDEKYSIEPSWECKFDHIEVRDGPF

447 DNGHIQSPNYPDDYRPSKVCIWRIQVSEGFHVGLTF-QSF BMP-1

188 GFSPIIGRFCGQQNPPVIKSSGRFLWIKFFADGELESMGFSARY BPC-1 506 ESSNLIGRYCGYENPDDIKSTSSRLWLKFVSDGSINKAGFAVNF BMP-1 * *** ** ** *** ** **

36.3% identity in 102 residues overlap in CUB domain 3 of BMP-1

130 GIFTSPNYPSKYPPDRECIYIIEAAPRQCIELYFDEKYSIEPSWECKFDHIEVRDGPFGF BPC-1 605 GSITSPGWPKEYPPNKNCIWQLVAPTQYRISLQFDF-FETEGNDVCKYDFVEVRSGLTAD BMP-1 *** * ***

190 SPIIGRFCGQQNPPVIKSSGRFLWIKFFADGELESMGFSARY BPC-1 664 SKLHGKFCGSEKPEVITSQYNNMRVEFKSDNTVSKKGFKAHF BMP-1 * *** * ** *

37.9% identity in 103 residues overlap in CUB domain 4 of BMP-1

130 GIFTSPNYPSKYPPDRECIYIIEAAPRQCIELYFDEKYSIEPSWECKFDHIEVRDGPFGF BPC-1 761 GTITSPNWPDKYPSKKECTWAISSTPGHRVKLTFVEM-DIESQPECAYDHLEVFDGRDAK BMP-1 * * * * * ** ** ** ** **

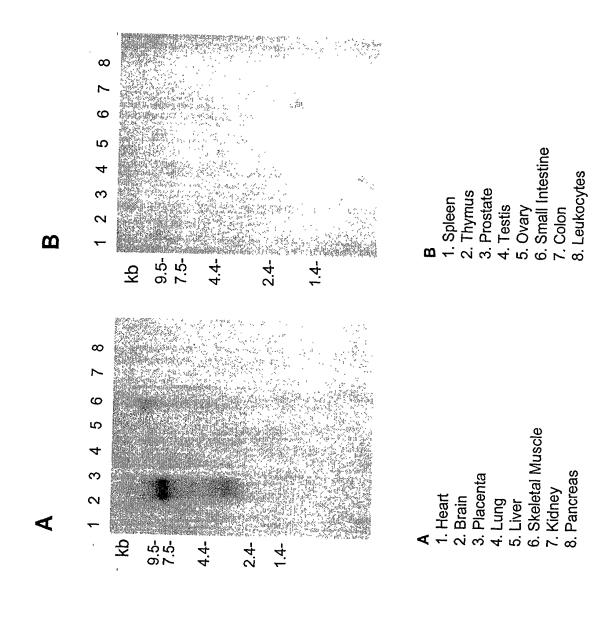
190 SPIIGRFCGQQNPPVIKSSGRFLWIKFFADGELESMGFSARYN BPC-1 820 APVLGRFCGSKKPEPVLATGNRMFLRFYSDNSVQRKGFQASHS BMP-1

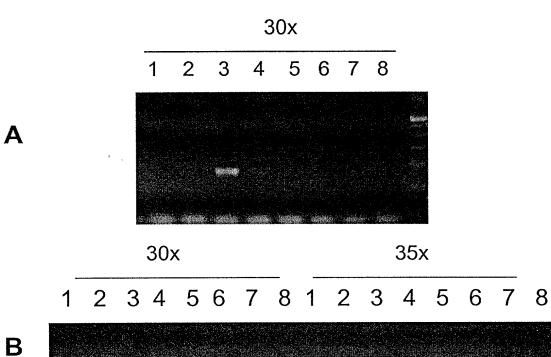
32.3% identity in 96 residues overlap in CUB domain 5 of BMP-1

139 SKYPPDRECIYIIEAAPRQCIELYFDEKYSIEPSWECKFDHIEVRDGPFGFSPIIGRFCG BPC-1 887 NNYPGGVDCEWVIVAEEGYGVELVF-QTFEVEEETDCGYDYIELFDGYDSTAPRLGRYCG BMP-1 ** * * * * * * * * * **

199 QQNPPVIKSSGRFLWIKFFADGELESMGFSARYNFT BPC-1 946 SGPPEEVYSAGDSVLVKFHSDDTISKKGFHLRYTST BMP-1

FIG. 7







Panels:

A

C

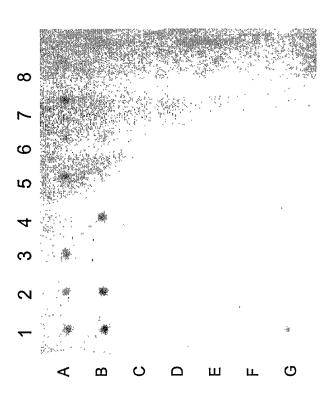
- 1. Brain
- 2. Prostate
- 3. LAPC-4 AD
- 4. LAPC-4 Al
- 5. LAPC-9 AD
- 6. HeLa
- 7. Murine cDNA
- 8. Neg. control

B

- 1. Brain
- 2. Heart
- 3. Kidney
- o. Idano
- 4. Liver
- 5. Lung
- 6. Pancreas
- 7. Placenta
- 8. Skeletal Muscle

C

- 1. Colon
- 2. Ovary
- 3. Leukocytes
- 4. Prostate
- 5. Small Intestine
- 6. Spleen
- 7. Testis
- 8. Thymus

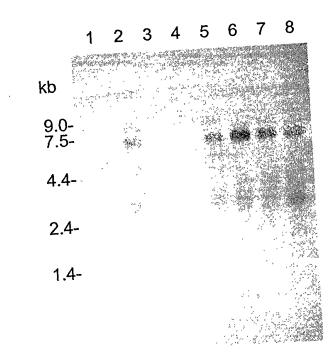


D1 testis D2 ovary D3 pancreas D4 pituitary gland D5 adrenal gland D6 thyroid gland D7 salivary gland D8 mammary gland	E1 kidney E2 liver E3 small intestine E4 spleen E5 thymus E6 peripheral leukocytes E7 lymph node E8 bone marrow
A1 brain A2 amygdala A3 caudate nucleus A4 cerebellum A5 cerebral cortex A6 frontal lobe A7 hippocampus A8 medulla oblongata	B1 occipital lobe B2 putamen B3 substantia nigra B4 temporal lobe B5 thalamus B6 sub-thalamic nucleus B7 spinal cord

G1 fetal brain
G2 fetal heart
G3 fetal kidney
G4 fetal liver
G5 fetal spleen
G6 fetal thymus
G7 fetal lung

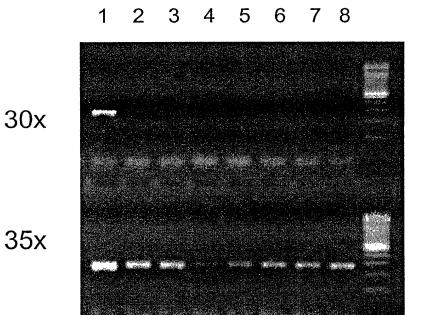
F1 appendix F2 lung F3 trachea F4 placenta

C2 aorta
C3 skeletal muscle
C4 colon
C5 bladder
C6 uterus
C7 prostate
C8 stomach



Lanes:

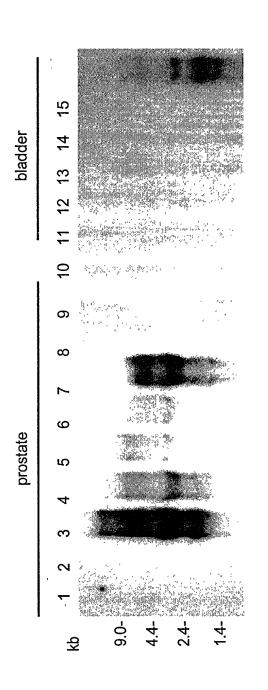
- 1. Cerebellum
- 2. Cerebral Cortex
- 3. Medulla
- 4. Spinal Cord
 5. Occipital Lobe
 6. Frontal Lobe
- 7. Temporal Lobe 8. Putamen



Panel:

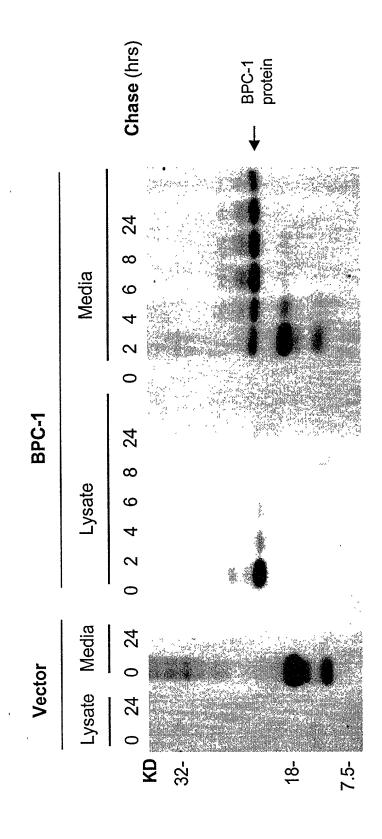
- 1. Brain
- 2. Heart
- 3. Kidney
- 4. Liver
- 5. Lung
- 6. Skeletal Muscle 7. Spleen 8. Thymus

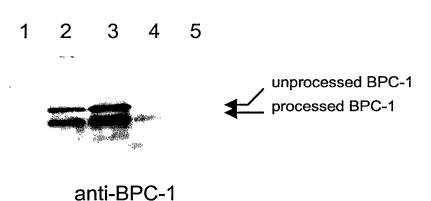
FIG. 9



1. PrEC 10. HT1197
2. Prostate 11. SCaBER
3. LAPC-4 AD 12. UM-UC-3
4. LAPC-4 AI 13. TCCSUP
5. LAPC-9 AD 14. J82
6. LAPC-9 AI 15. 5637
7. LNCaP 8. PC-3
9. DU145

FIG. 10

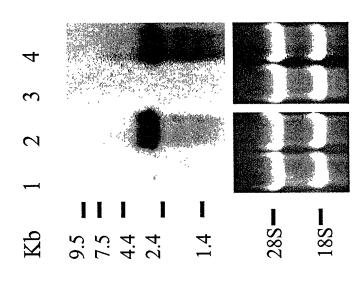




Lanes:

- 1. HighFive cell extract- uninfected
- 2. HighFive cell extract- infected with BPC-1 virus
- 3. HighFive cell extract infected with BPC-1 virus
- 4. HighFive cell media infected with BPC-1 virus
- 5. HighFive cell media infected with BPC-1 virus

FIG. 12



1. PC3/neo

2. PC3/BPC-1
3. 3T3CL7/neo, acute infection
4. 3T3CL7/BPC-1, acute infection

FIG. 13

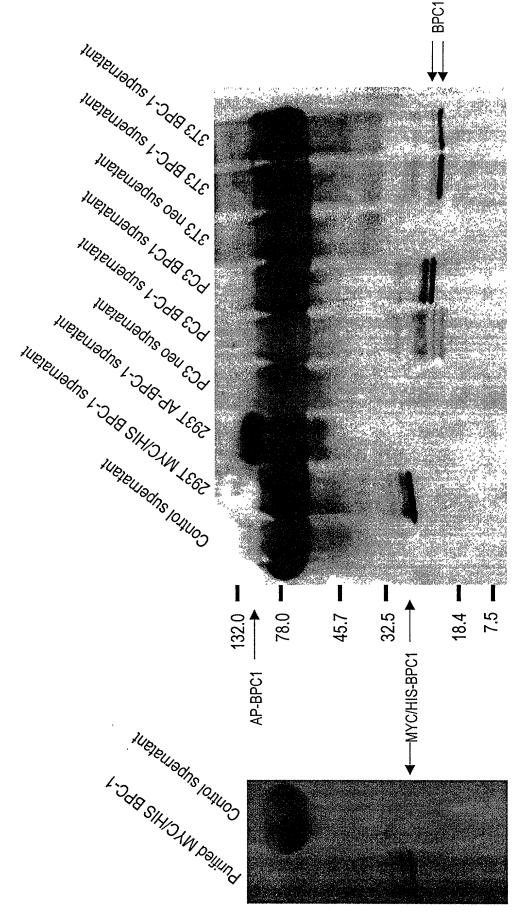
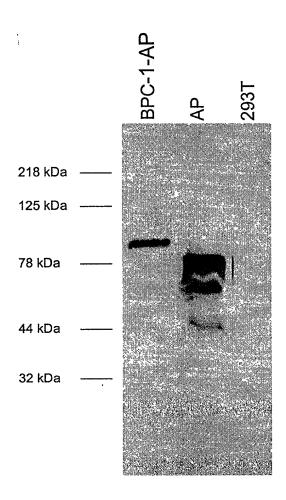


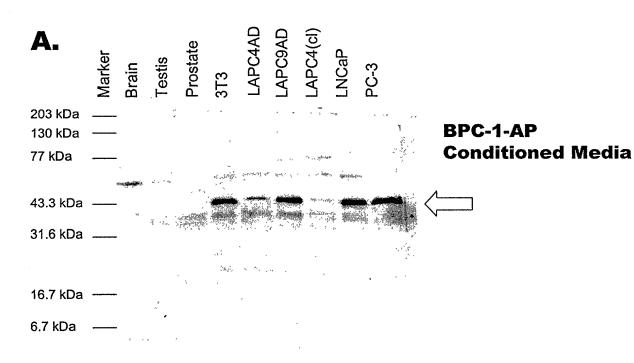
FIG. 14

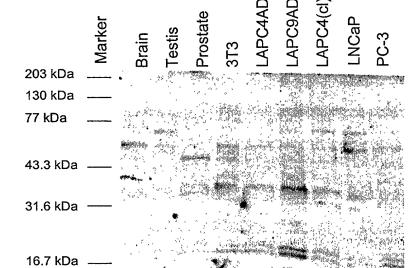


Anti-HIS

B.

6.7 kDa





Alkaline Phosphatase Conditioned Media

My residence, post office address and citizenship are as stated next to my name.

I believe I am the original, first and sole inventor (if only one name is listed below) or an original, first and joint inventor (if plural names are listed below) of the subject matter which is claimed and for which a patent is sought on the invention entitled:

DECLARATION AND POWER OF ATTORNEY

BPC-1: A SECRETED BRAIN-SPECIFIC PROTEIN EXPRESSED AND SECRETED BY PROSTATE AND BLADDER CANCER CELLS

the specification of which

was filed on August 10, 1999 as United States Application Number 09/374,135

I hereby state that I have reviewed and understand the contents of the above-identified specification, including the claims, as amended by any amendment specifically referred to above.

I acknowledge the duty to disclose information which is material to patentability as defined in and in accordance with the provisions of Title 37 of the Code of Federal Regulations, Section 1.56.

I hereby claim the benefit of under Title 35 of the United States Code, Section 119(e) of any United States provisional application(s) listed below.

Application Number

Filing Date

60/095,982

August 10, 1998

I hereby appoint:

Kenneth K. Sharples

Reg. No. 35,355

as my attorney(s) and/or patent agent(s), with full power of substitution and revocation, to prosecute this application and to transact all business in the Patent and Trademark Office in connection therewith.

I further direct that all correspondence concerning this application be directed to:

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Point Reyes Station, CA 94956

I hereby declare that all statements made herein of my own knowledge are true and that all statements made on information and belief are believed to be true; and further that these statements were made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United States Code and that such willful false statements may jeopardize the validity of the application or any patent issued thereon.

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